

http://es/ScoreAccessWeb/GetItem.action?AppId=10063516&seqId=631147&ItemName=us-... 9/8/06

SCORE Search Results Details for Application 10063516 and Search Result us-10-063-516-12.rapbn.

Score Home	Retrieve Application	SCORE System	SCORE	Comments /
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This page gives you Search Results detail for the Application 10063516 and Search Result us-10-063-516-12.rapbn.

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GenCore version 5.1.9
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OM protein - protein search, using sw model

Run on: September 6, 2006, 16:58:37 ; Search time 33 Seconds
(without alignments)
972.853 Million cell updates/sec

Title: US-10-063-516-12
Perfect score: 2448
Sequence: 1 MGACLGACSLLSASCSCGSG.....LLLYLWTLVAPLLLRNRDFS 457

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 254368 seqs, 70249769 residues

Total number of hits satisfying chosen parameters: 254368

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_AA_New:*
1: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_NEW_PUB.pep:*
5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
6: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
7: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US11_NEW_PUB.pep:*
8: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US60_NEW_PUB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result	% Query
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OM protein - protein search, using sw model

Run on: September 6, 2006, 16:47:36 ; Search time 46 Seconds
(without alignments)
997.505 Million cell updates/sec

Title: US-10-063-516-12
Perfect score: 2448
Sequence: 1 MGACLGACSLLSASCSCGSG.....LLLYLWTLVAPLLLRNRDFS 457

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 409781 seqs, 100405355 residues

Total number of hits satisfying chosen parameters: 409781

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Pending Patents_AA_New:*

- 1: /EMC_Celerra_SIDS3/ptodata/2/paa/PCT_NEW_COMB.pep:*
- 2: /EMC_Celerra_SIDS3/ptodata/2/paa/US06_NEW_COMB.pep:*
- 3: /EMC_Celerra_SIDS3/ptodata/2/paa/US07_NEW_COMB.pep:*
- 4: /EMC_Celerra_SIDS3/ptodata/2/paa/US08_NEW_COMB.pep:*
- 5: /EMC_Celerra_SIDS3/ptodata/2/paa/US09_NEW_COMB.pep:*
- 6: /EMC_Celerra_SIDS3/ptodata/2/paa/US10_NEW_COMB.pep:*
- 7: /EMC_Celerra_SIDS3/ptodata/2/paa/US11_NEW_COMB.pep:*
- 8: /EMC_Celerra_SIDS3/ptodata/2/paa/US60_NEW_COMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	%
Result	Query